

# Pattern discovery for semi-structured web pages using bar-tree representation

Z. Akbar, L.T. Handoko

**Abstract**—Many websites with an underlying database containing structured data provide the richest and most dense source of information relevant for topical data integration. The real data integration requires sustainable and reliable pattern discovery to enable accurate content retrieval and to recognize pattern changes from time to time; yet, extracting the structured data from web documents is still lacking from its accuracy. This paper proposes the bar-tree representation to describe the whole pattern of web pages in an efficient way based on the reverse algorithm. While previous algorithms always trace the pattern and extract the region of interest from *top root*, the reverse algorithm recognizes the pattern from the region of interest to both top and bottom roots simultaneously. The attributes are then extracted and labeled reversely from the region of interest of targeted contents. Since using conventional representations for the algorithm should require more computational power, the bar-tree method is developed to represent the generated patterns using bar graphs characterized by the depths and widths from the document roots. We show that this representation is suitable for extracting the data from the semi-structured web sources, and for detecting the template changes of targeted pages. The experimental results show perfect recognition rate for template changes in several web targets.

**Keywords**—data extraction, data mining, web-based information system

## I. INTRODUCTION

Text mining, especially from the web sources is getting important during the last decade. This is triggered by the exponentially growing number of websites with various types of information on the net. Most of them are providing the information generated from the structured data in an underlying database through certain predefined templates or layouts [1]. Following the great number of web pages in this kind which are already available on the net, these semi-structured web sources contain rich and unlimited valuable data for a variety of purposes. Extracting those data and then rebuilding them into a structured database are a challenge to realize an automatic data mining from web sources.

Several methods for these purposes have been proposed previously in the literature. Some of them can be classified as the so-called wrappers [2], [3], [4], [5], [6] which have been briefly surveyed in [7]. The wrapper technique allows an automatic data extraction through predefined wrapper created for each target data source. The wrappers then accepts a query against the data source and returns a set of structured results to the calling application.

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On the other hand, there are several automatic methods without a manual initial learning process. For example, some methods are generating the template automatically from first multiple pages before extracting the rest of data based on the template [8], [9], [10]. A more comprehensive method without requiring multiple pages has also been proposed using a page creation model which captures the main characteristics of semi-structured web pages to derive the set of properties [11]. Though, the last method is more intended to extract the lists of data records from a single web page with sibling subtrees.

Since 2008 our group has worked on developing an online infrastructure with a major purpose of integrating the information related to science and technology across Indonesia, that is the Indonesian Scientific Index – ISI [12]. However, in contrast with conventional approach where the data are collected through official connection under certain regulation, ISI integrates the data indirectly by harvesting certain web contents of the official websites of targeted institutions. Initially the method was motivated by the failures of some *conventional* methods of data integration which always requires certain standard at any level and leads to additional works in the participating institutions. On the other hand, as a part of public responsibility all academic institutions have developed and launched various public information through their own websites. Therefore the idea of indirect data integration is welcomed by all participating institutions, since it is not like a dictatorship, more acceptable, much cheaper and more efficient for all parties than the conventional one which requires a kind of standardization among the information islands belong to separated institutions [13], [14]. The main problem is yet improving the accuracy of data retrieval and restructuring them into desired fields for further content analysis.

The architecture of ISI is inspired and the combination of focused web-crawling and regular web-harvesting. The focused web-crawling does not indiscriminately crawl the web pages like general purpose search engines, but attempts to download pages that are similar to each other [15]. Here, we follow the same line to harvest certain types of web pages with specific contents. Throughout the paper, let us call this method as the focused web-harvesting [14]. Nonetheless, as an official data integrator it must not allow any errors in the data collection. According to the data integration purposes and its requirement of high accuracy, the first version of focused web-harvesting technique at ISI adopts human intervention in the initial setup by providing the targeted URL of list of data, for instance the publication list, and defining the template of the final page contains the relevant information by labeling the attributes. In the case of detail information of a publication

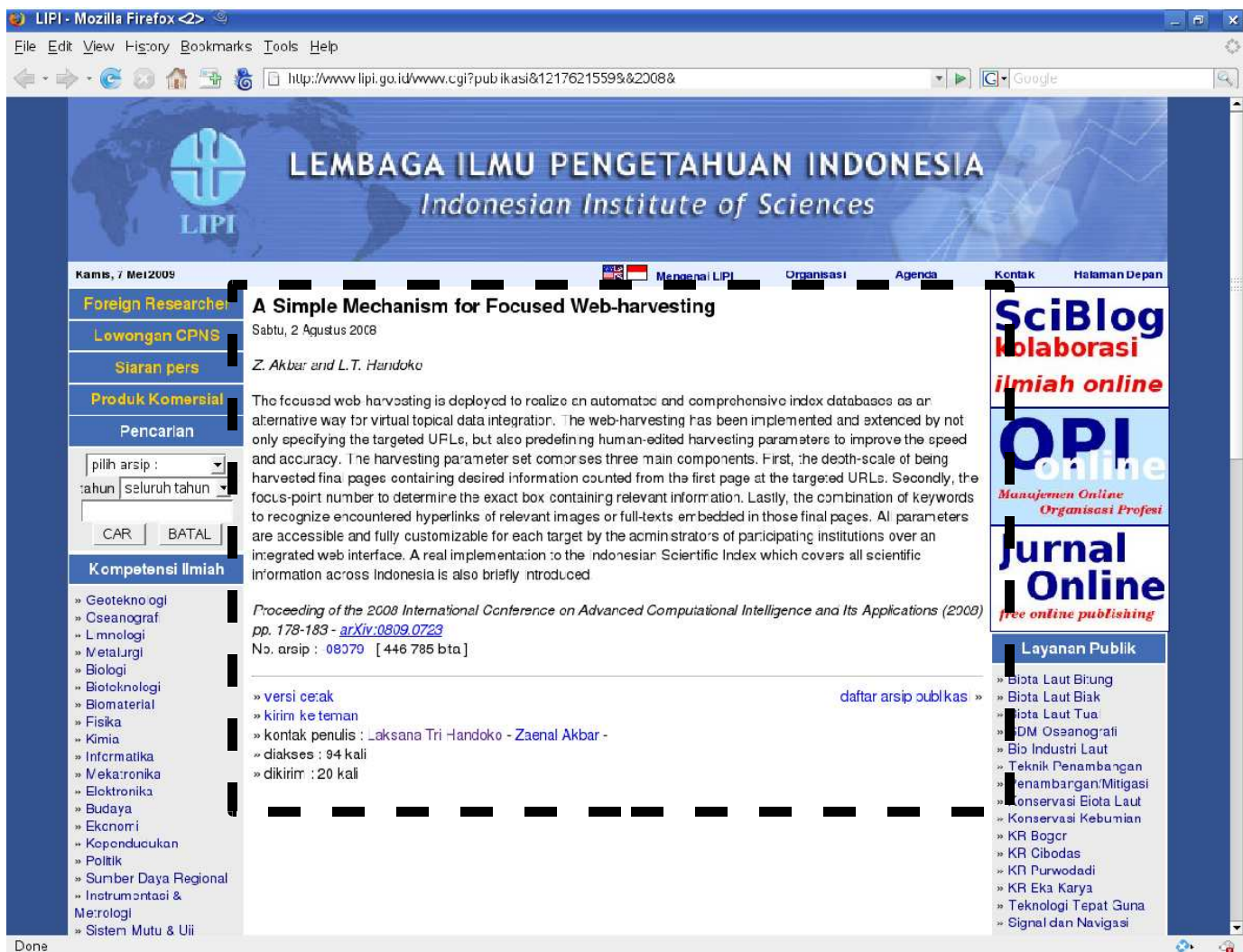


Fig. 1. The example of the desired RoI from the displayed content of a scientific publication page on the web, shown by the areas inside the dashed rectangular.

page as shown in Figs. 1 and 2, the relevant information and labeling are ranging from title, authors, abstract to the full-text if available. Comparing with the general purpose crawling [16], or even focused web-crawling [17], obviously ISI could retrieve the information more accurately due to its targeted contents and sources. The approach has also been realized and integrated in a user friendly web-based interface to enable the administrator to set up the initial parameters for each targeted sources. The toolkit has been released as an open source under GNU Public License at SourceForge.net [18].

Unfortunately, in spite of its high accuracy and unnecessary machine-learning like system, the method is suffered from tedious labor time at its initial setup to determine the region of interest (RoI), the tokens and to label the attributes. Moreover, it is lacking of the ability to detect effectively later changes of the targeted page templates. Any later changes of the targeted templates will again require human intervention to manually revise the parameters accordingly. Especially, because the labels and tokens are represented as DOM trees which are sensitive to later changes of targeted web templates [10], [19].

On the other hand, we have also found that the previously known methods like wrapper induction [3], or the information extraction based on multi pattern discovery techniques [20], are not suitable for our purpose since all of them by definition contain significant statistical errors which could burdens the initial purpose of official data integration.

In our recent work [21], in order to overcome the above-mentioned problems a new method has been introduced to extract the RoI at the final targeted web pages and its tokens reversely from the RoI to both top and bottom roots, and further to label each attribute as usual. The technique is in contrast with the existing mechanisms which always start from the top root of targeted web pages or its RoI. It has been argued that the so-called reverse algorithm is more efficient and accurate to extract the tokens, and on the other hand to detect later template changes without any ambiguities. We should also remark that the algorithm is applicable for any existing methods for data extraction, especially the ones which require initial setup by human intervention to define and label the attributes. This is actually similar to the previous method



- 1) Determine the URL address of the final web page with desired information like Fig. 1.
- 2) Provide the whole sentences of the RoI by copying and pasting the 'desired text' displayed on screen, not its source.
- 3) Provide the whole sentence(s) of each sub-RoI and assign the attributes for each of them.
- 4) Crawl the source of the final web page at 1.
- 5) Parse and clean the text-format HTML tags like `<b>`, `<i>`, etc.
- 6) Take the upper part of source from the top till the last one before the first sentence of RoI. Parse and clean all texts inside except the layout-format HTML tags, like `<tr>`, `<span>`, etc. Do the same thing for the lower part that is from the end of last sentence of RoI till the bottom.
- 7) Count the number of 'open-tag' ( $n_{ot}$ ) and 'closed-tag' ( $n_{ct}$ ) from the deepest part in term of desired content, that is the nearest tags from the RoI.

We should stress here that there is no need for the administrators to provide the web page sources at all. Open-tag here means the tags which have no pair in upper or lower part, while the closed-tag denotes the pairing tags within upper or lower part. Of course, our interest is only in the open-tag which should describe the whole structure of web template.

Following the above procedures, we can obtain a kind of DOM tree as shown in Fig. 3. We can further count the number of trees according to the number of open-tags. Please remind that the counting is done horizontally, from left to right shown by the arrow in the figure. The number of trees in upper and lower parts are determined by,

$$\Sigma \equiv n_{ot} - n_{ct} . \quad (1)$$

Concerning all possibilities on the number of trees in upper and lower parts, therefore we can generally categorize the web structures through the discrepancies between both numbers as,

$$\Delta \equiv \Sigma_{upper} - \Sigma_{lower} \quad (2)$$

$$\begin{cases} = 0 & : \text{fully symmetry} \\ < 0 & : \text{lower asymmetry} \\ > 0 & : \text{upper asymmetry} \end{cases}$$

Fig. 3 provides an example of tree in the case of Fig. 1 which is accidentally asymmetry. That means the number of trees in the upper and lower parts are not the same,  $\Sigma_{upper} \neq \Sigma_{lower}$ . Again, we can use one of the existing methods to calculate the number of trees like the PAT tree algorithm [22] and so forth.

Through the discussion above, it is clear that the present method has several advantages :

- We can separate independently the structure and the rules to obtain the RoI and the structure inside.
- We can find out the template changes and its relevance with the desired RoI, since we can compare and see the pairing tokens between the upper and lower parts.
- There is no need for further human intervention as long as the page containing the initial RoI still exist. The system uses the same RoI as keyword to perform regular check

to detect template changes before recrawl the same target. Only if the content is removed by the owner, the system will defer the recrawling job at the target and send a warning to the administrator to choose another content as new keyword.

We discuss these points in more detail through the real implementation at ISI in the subsequent Sec. IV.

Further issue is then how to represent the method, not only visually, but also mathematically to enable more quantitative analysis in real implementation.

### III. THE BAR-TREE REPRESENTATION

Here we introduce the way to represent the reverse algorithm method in form of bar diagrams. The bar is characterized by its width ( $w$ ) and height. The height is determined by the depth ( $d$ ) of each column of attributes from the root document. On the other hand, the width is given by the number of parallel attributes ( $P_d$ ) in certain depth weighted by a ratio ( $r$ ) according to the depth. The definition is illustrated in Figs. 3 and 4.

Considering the mentioned-above definition, the width of each bar diagram can be written as follows,

$$w_d = \begin{cases} I & \text{for } d = 0 \\ \frac{I - (d-1)r}{P_{d-1}} w_{d-1} & \text{for } d > 0 \end{cases} , \quad (3)$$

where  $I$  is the given initial width and  $r \leq I/d_{\max}$  is the appropriate ratio to decrease the width following the depth of a bar. The parallel attributes  $P_d$  is nothing else than the number of attributes at the same  $n$ -th depth. For instance, the 5th column in Fig. 3 has  $P_5 = 4$ , while at the 9th column  $P_9 = 3$  and so forth.

According to the definition, the bar-tree representation in the case of Fig. 3 can be further depicted as Fig. 4. One should remark a rule that for  $P_d > 1$ , the  $(d+1)$ th bar should be drawn inside the appropriate order of bar in that depth. In the case of  $d = 5$  in Fig. 4, since  $P_5 = 4$  and the interested bar is the 3rd one, the  $d = 6$ 's bar is put inside the 3rd of  $d = 5$ 's bar accordingly.

Quantitatively, the pattern of bar-tree representation can be characterized by its partial and total squares. The square of each individual bar is given by,

$$A_d = \begin{cases} 0 & \text{for } d = 0 \\ d \times \frac{I - (d-1)r}{P_{d-1}} w_{d-1} & \text{for } d > 0 \end{cases} . \quad (4)$$

The "nett-square", that means the square of bar which is not overlapped with its lower bars, for each bar is given by,

$$\begin{aligned} A_d^{\text{nett}} &= d \times \frac{I - (d-1)r}{P_{d-1}} w_{d-1} - (d-1) \times \frac{I - (d-1)r}{P_{d-1}} w_{d-1} \\ &= \frac{I - (d-1)r}{P_{d-1}} w_{d-1} . \end{aligned} \quad (5)$$

Using Eqs. (3) and (4), it subsequently leads to,

$$A_d^{\text{nett}} = \left[ \prod_{n=0}^{d-1} \frac{I - (d-1-n)r}{P_{d-1-n}} \right] w_0 . \quad (6)$$





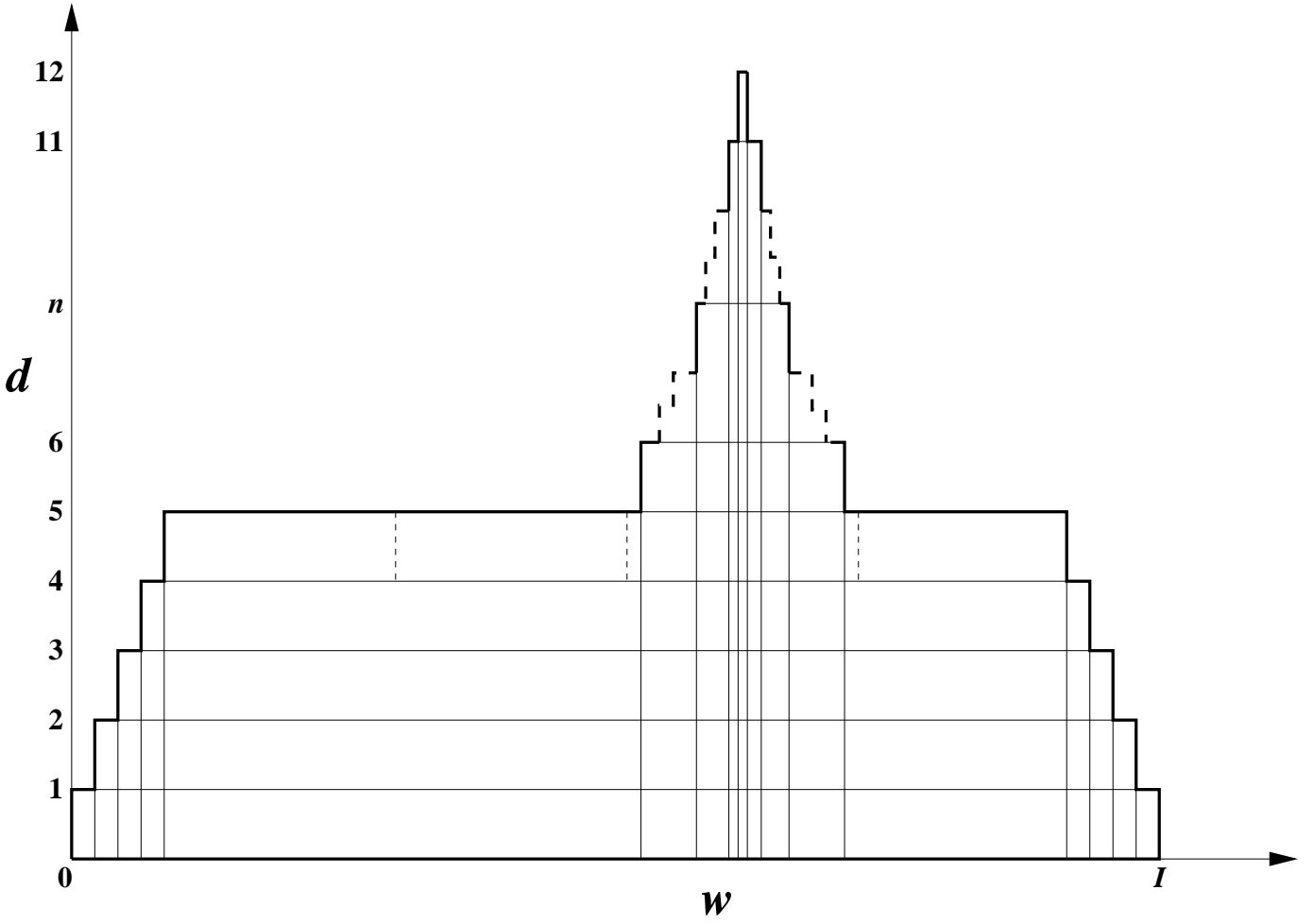


Fig. 4. The bar-tree representation with  $w$  and  $d$  denote the width and depth of each tree as depicted in Fig. 3 for  $r = 10\% \times I$ .

The results for two cases representing the calculation using different variable sets,  $\{d_{\max}, A^{\text{total}}\}$  and  $\{d_{\max}, A^{\text{total}}, P_d, A_d\}$ , are shown in Figs. 6 and 7. The time consumption in Fig. 6 means the running time in millisecond required for the whole processes from parsing the HTML till calculating the whole variables defined above. While the accuracy rate in Fig. 7 represents the successful rate to detect the template changes. The template changes were done randomly but intensively among the data using certain script. We should note that the template changes are completely same for both variable sets.

From the figures, we can deduce several points :

- As the number of  $d_{\max}$  is greater, it requires more time to calculate all variables and also decreases slightly the accuracy. The reason is obvious, since the trees with deeper structure have more probabilities and complexities of template changes.
- Using more complete variable set  $\{d_{\max}, A^{\text{total}}, P_d, A_d\}$  would improve significantly the accuracy to detect the template changes than the simpler one  $\{d_{\max}, A^{\text{total}}\}$  without significant increasing in time. Because the variables  $A^{\text{total}}$  could be accidentally the same if the template changes occurred at the same depth  $d$ . For instance, if only the sequence in a depth  $d$  is different,

then  $P_d$  should remain unaltered.

- Although the variable set  $\{d_{\max}, A^{\text{total}}, P_d, A_d\}$  is enough in most applications, yet there is no guarantee to correctly detect the location of template changes. Fortunately, it is indeed not necessary in the reverse algorithm. Because once the template changes were detected, the new pattern is re-extracted from the RoI to replace the old one, no matter where the changes happened.

If one requires, for certain needs, more accurate detection power, that is 100% in our experimental running, we recommend to perform further check using the variable  $\Delta$  in Eq. (2). The discrepancies between the old and new numbers of  $\Delta$  would unambiguously detect the template changes and its exact location. It can be summarized as below,

- 1) No change at all :  
 $\Delta^{\text{new}} = \Delta^{\text{old}}, \Sigma_{\text{upper}}^{\text{new}} = \Sigma_{\text{upper}}^{\text{old}}, \Sigma_{\text{lower}}^{\text{new}} = \Sigma_{\text{lower}}^{\text{old}}$
- 2) Simultaneous changes with same size in both upper and lower trees :  
 $\Delta^{\text{new}} = \Delta^{\text{old}}, \Sigma_{\text{upper}}^{\text{new}} \neq \Sigma_{\text{upper}}^{\text{old}}, \Sigma_{\text{lower}}^{\text{new}} \neq \Sigma_{\text{lower}}^{\text{old}}$
- 3) Only either upper or lower tree has changed :  
 $\Delta^{\text{new}} \neq \Delta^{\text{old}}, \Sigma_{\text{upper}}^{\text{new}} \neq \Sigma_{\text{upper}}^{\text{old}}, \Sigma_{\text{lower}}^{\text{new}} = \Sigma_{\text{lower}}^{\text{old}}$   
or :  
 $\Delta^{\text{new}} \neq \Delta^{\text{old}}, \Sigma_{\text{upper}}^{\text{new}} = \Sigma_{\text{upper}}^{\text{old}}, \Sigma_{\text{lower}}^{\text{new}} \neq \Sigma_{\text{lower}}^{\text{old}}$

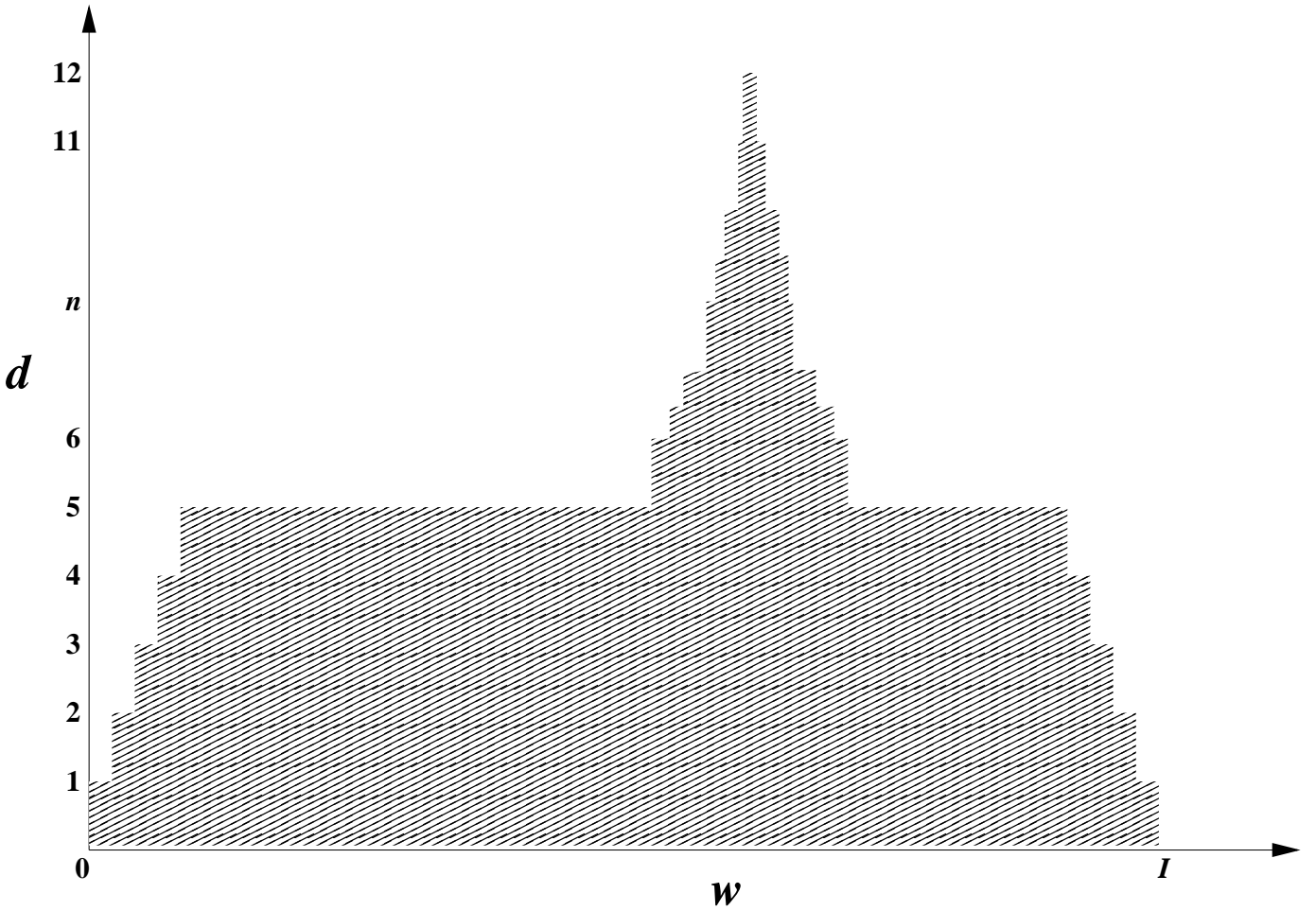


Fig. 5. The total pattern of bar-tree representation in the case of Figs. 3 and 4.

- 4) Both upper and lower trees have changed differently :  
 $\Delta^{\text{new}} \neq \Delta^{\text{old}}; \Sigma_{\text{upper}}^{\text{new}} \neq \Sigma_{\text{upper}}^{\text{old}}; \Sigma_{\text{lower}}^{\text{new}} \neq \Sigma_{\text{lower}}^{\text{old}}$

Apparently, in the case 1 no need to alter the stored initial variables. In contrast, from the case 2, 3 and 4 we can deduce that the templates have been changed, either in the upper, lower or both trees.

## V. CONCLUSION

We have discussed the bar-tree representation suitable for the reverse algorithm method to extract the RoI and to label the relevant attributes in the initial setup. The resulted patterns can be used further to automatically extract the data from crawled targeted web pages. We argue that the method with additional relevant web interface would reduce the administrator works significantly. On the other hand it improves the accuracy and speed of finding the tokens and labeling the attributes. Because the human intervention is basically required only during the initial setup. The pattern recognition in this method is done in an exact way without, for instance, any predefined parameters like threshold value etc.

We have found that this method is very effective to detect the template changes, for instance newly inserted banners in the middle of upper or lower tree which often occurs in any

websites and leads to difficulties in existing methods. The important point is it does not require huge computer power, nor further human intervention once the initial setup has been done.

Through our experimental running, we can conclude that the focused web-harvesting deploying the combination of reverse algorithm and bar-tree representation is appropriate for the indirect data integration. The method performs the data collection over targeted web sources very accurately. We also recommend to perform the full set  $\{d_{\text{max}}, A^{\text{total}}, P_d, A_d\}$  rather than the simpler set  $\{d_{\text{max}}, A^{\text{total}}\}$  to obtain results with much higher accuracy in a moderate time consumption.

The real works on applying the method to further restructure the huge number of data stored at ISI is still under progress. The results and its effectiveness to restructure all relevant fields in a nation wide scientific index will be analyzed and reported elsewhere. Finally, we should remark that the method is also applicable for the web sources in a form of list of data. The work on this matter is also in progress.

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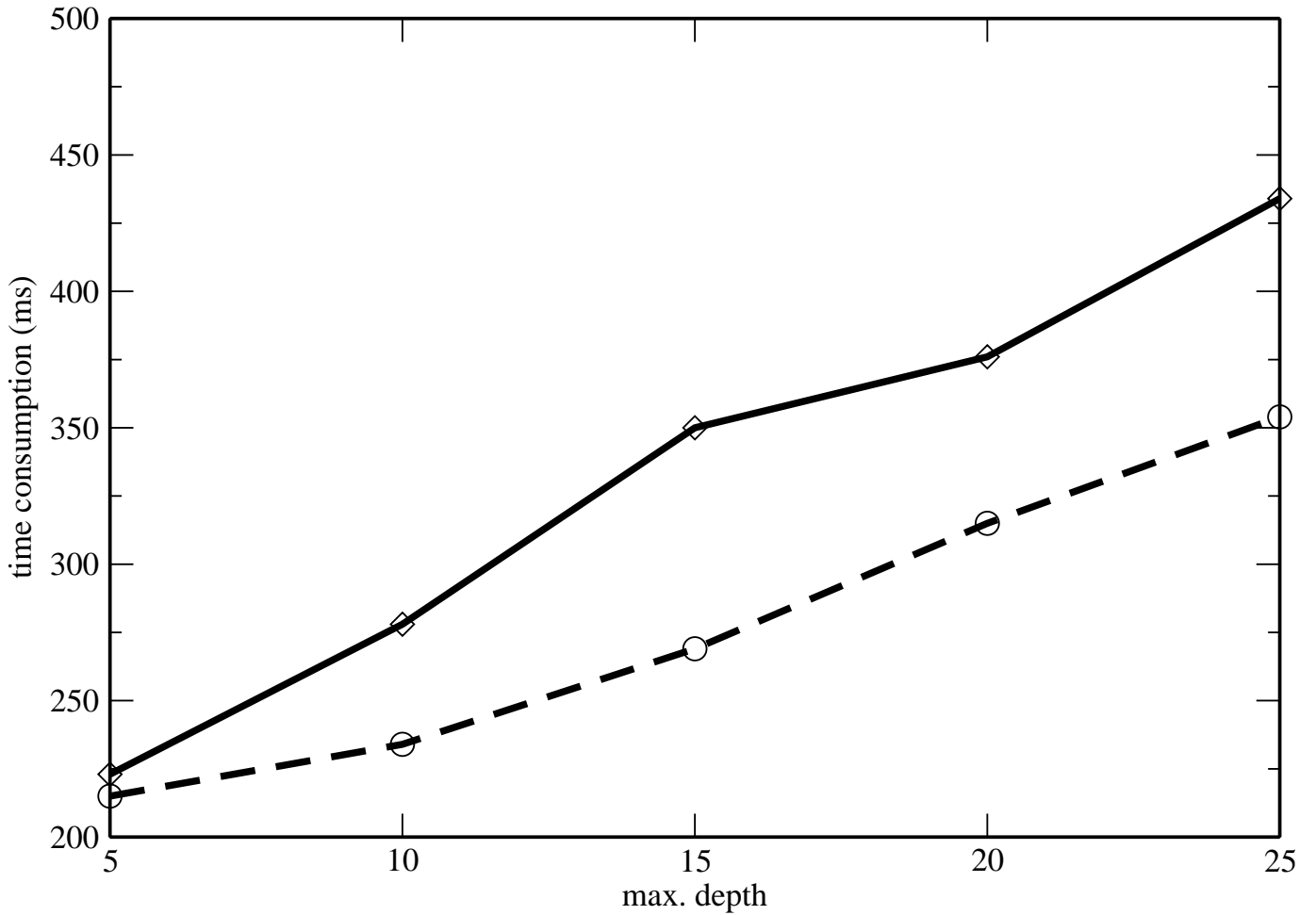


Fig. 6. The measurement result of time consumption for detecting the template changes with various number of  $d_{\max}$ . The solid and dashed lines denote the performance with variable sets :  $\{d_{\max}, A^{\text{total}}, P_d, A_d\}$  and  $\{d_{\max}, A^{\text{total}}\}$ .

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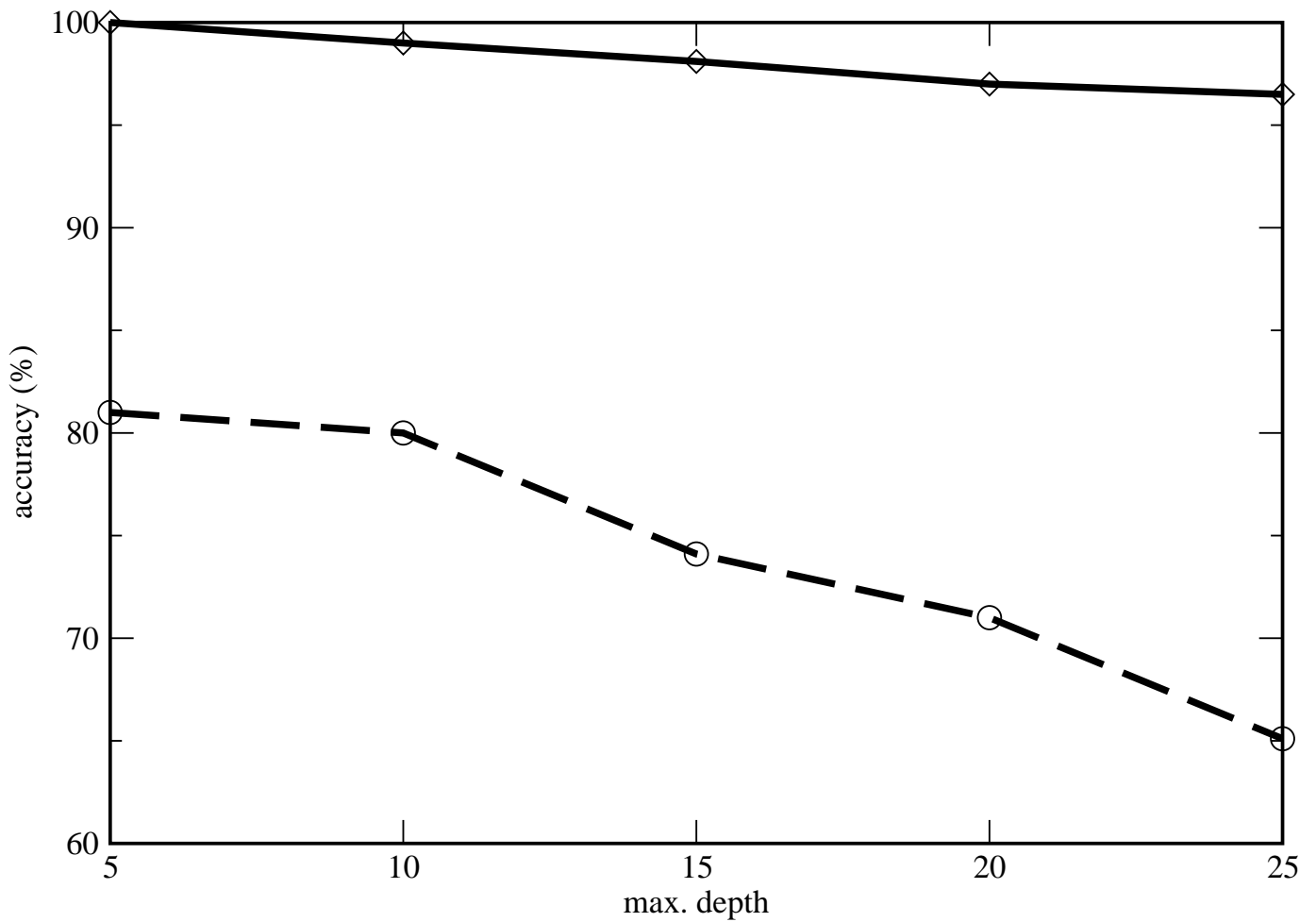


Fig. 7. The measurement result of accuracy for detecting the template changes with various number of  $d_{\max}$ . The solid and dashed lines denote the performance with variable sets :  $\{d_{\max}, A^{\text{total}}, P_d, A_d\}$  and  $\{d_{\max}, A^{\text{total}}\}$ .

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